Amendments to the Claims

- 1. (Currently Amended) A method for identifying a <u>candidate</u> modulatory compound that is <u>capable of for</u> decreasing the expression or activity of a *daf-16* gene, said method comprising:
 - (a) providing a <u>C. elegans or nematode</u>, isolated <u>C. elegans nematode</u> cell , or isolated mammalian cell expressing a daf-16 gene; and
- (b) contacting said a <u>C. elegans</u> or nematode, isolated <u>C. elegans</u> nematode cell, or isolated mammalian cell with a candidate compound, wherein a decrease in daf-16 expression or activity following contact of said <u>C. elegans</u> nematode, or said isolated <u>C. elegans</u> nematode cell, or said isolated mammalian cell with said candidate compound identifying identifies a candidate modulatory compound.
- 2. (Original) The method of claim 1, wherein said compound is a candidate compound for ameliorating or delaying an impaired glucose tolerance condition, atherosclerosis, or obesity.
 - 3. (Cancelled)
- 4. (Original) The method of claim 1, wherein said *daf-16* gene is a nematode *daf-16* gene.

- 5. (Withdrawn) A method for the identification of a compound that is a candidate compound for ameliorating or delaying an impaired glucose tolerance condition, said method comprising the steps of:
 - (a) providing a daf-2, daf-16 mutant nematode;
- (b) expressing in the cells of said nematode a mammalian AFX polypeptide, whereby said nematode forms a dauer larva; and
- (c) contacting said dauer larva with a compound, wherein release from the dauer larval state is an indication that said compound is a candidate compound for ameliorating or delaying an impaired glucose intolerance condition.
- 6. (Withdrawn) A method for the identification of a compound that is a candidate compound for ameliorating or delaying an impaired glucose tolerance condition, said method comprising the steps of:
 - (a) providing an age-1, daf-16 mutant nematode;
- (b) expressing in the cells of said nematode a mammalian AFX polypeptide, whereby said nematode forms a dauer larva; and
- (c) contacting said dauer larva with a compound, wherein release from the dauer larval state is an indication that said compound is a candidate compound for ameliorating or delaying an impaired glucose intolerance condition.

- 7. (Withdrawn) A method for the identification of a compound that is a candidate compound for ameliorating or delaying an impaired glucose tolerance condition, said method comprising the steps of:
 - (a) providing a daf-2, daf-16 mutant nematode;
- (b) expressing in the cells of said nematode a mammalian FKHR polypeptide, whereby said nematode forms a dauer larva; and
- (c) contacting said dauer larva with a compound, wherein release from the dauer larval state is an indication that said compound is a candidate compound for ameliorating or delaying an impaired glucose intolerance condition.
- 8. (Withdrawn) A method for the identification of a compound that is a candidate compound for ameliorating or delaying an impaired glucose tolerance condition, said method comprising the steps of:
 - (a) providing an age-1, daf-16 mutant nematode;
- (b) expressing in the cells of said nematode a mammalian FKHR polypeptide, whereby said nematode forms a dauer larva; and
- (c) contacting said dauer larva with a compound, wherein release from the dauer larval state is an indication that said compound is a candidate compound for ameliorating or delaying an impaired glucose intolerance condition.

- 9. (Withdrawn) The method of any of claims 5-8, wherein said nematode is *C. elegans*.
- 10. (Withdrawn) The method of any of claims 5-8, wherein said compound is a candidate compound for ameliorating or delaying an impaired glucose tolerance condition that involves obesity or atherosclerosis.
- 11. (Withdrawn) A method for identifying a compound that modulates the interaction between DAF-16 and a second DAF polypeptide, said method comprising the steps of:
 - (a) providing a DAF-16 polypeptide;
 - (b) providing a second DAF polypeptide;
- (c) allowing said DAF-16 polypeptide and said second DAF polypeptide to interact and form a complex;
- (c) contacting said complex with a candidate compound, a modulation in the interaction between said DAF-16 and said second DAF polypeptide identifying a modulatory compound.
 - 12. (New) The method of claim 1, wherein said daf-16 gene has at least 71%

amino acid sequence identity to SEQ ID NO:54.

- 13. (New) The method of claim 1, wherein said daf-16 gene is a human gene.
- 14. (New) The method of claim 13, wherein said human gene is AFX.
- 15. (New) The method of claim 13, wherein said human gene is FKHR.